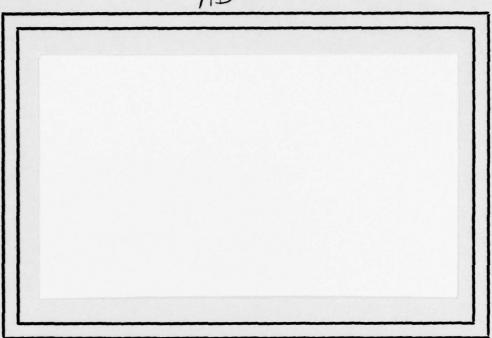


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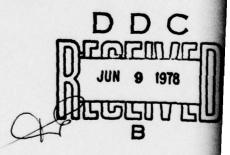
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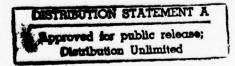
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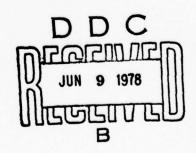
CELLULAR GRAPH ACCEPTORS, 3

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#### **ABSTRACT**

In earlier reports, cellular acceptors were studied whose languages are sets of d-graphs, i.e., labelled graphs of bounded degree whose arcs at each node are numbered. This report discusses acceptance tasks that depend on the concept of d-graph isomorphism -- in particular, the task of deciding whether a d-graph has a d-subgraph isomorphic to a given d-graph.





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#### 1. Introduction

Cellular acceptors whose languages are sets of d-graphs (labelled graphs of bounded degree whose arcs at each node are numbered) were studied in [1-2], where the terminology and notation used in the present paper are defined. This paper discusses acceptance tasks that depend on the concept of d-graph isomorphism -- in particular, the task of deciding whether a d-graph has a d-subgraph isomorphic to a given d-graph.

Given two node labelled graphs  $\gamma_1 = (N_1, A_1, f_1)$  and  $\gamma_2 = (N_2, A_2, f_2)$  where  $f_1, f_2$  are the node labelling functions,  $\gamma_1$  is <u>isomorphic</u> to  $\gamma_2$  if there exists a bijection b from  $N_1$  to  $N_2$  such that  $f_1(n) = f_2(b(n)) \forall n \in N_1$  and  $(m,n) \in A_1$ iff.  $(b(m), b(n)) \in A_2$ . A  $d_1$ -graph  $\Gamma_1 = (N_1, A_1, f_1, g_1)$  and a  $d_2$ -graph  $\Gamma_2 = (N_2, A_2, f_2, g_2)$  are <u>isomorphic</u> (denoted by  $\Gamma_1 \simeq \Gamma_2$ ) iff. their underlying graphs  $U(\Gamma_1)$  and  $U(\Gamma_2)$  are isomorphic. Here we allow  $d_1 \neq d_2$ . A <u>subgraph</u> of a d-graph  $\Gamma$ = (N, A, f, g) is denoted by (N', A', f|N', g|A') where N'⊆N and A'⊆A and if (m,n) tA' then mtN' and ntN'. Note that (N', A', f | N', g | A') is not necessarily a d-graph, since some of the nodes may not have exactly d neighbors. However, we can always attach # nodes so as to make it into a d-graph. A labelled graph  $\alpha$  is isomorphic to  $\Gamma$  if  $\alpha = U(\Gamma)$ , and  $\alpha$  is isomorphic to a subgraph of  $\Gamma$  if  $\alpha \simeq U(\Gamma')$  for some subgraph  $\Gamma'$  of  $\Gamma$ . In the following, we will consider only connected d-graphs. 

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#### 2. Graph isomorphism

In this section, we will consider acceptance tasks that depend on graph isomorphism. Specifically, given a labelled graph  $\alpha$  of degree  $\leq$  d, we will find a finite state acceptor  $\mathbf{M}_{\alpha}$  such that  $(\Gamma, \mathbf{M}_{\alpha}, \mathbf{H})$  accepts  $\Gamma$  iff.  $\alpha$  is isomorphic to  $\Gamma$  and rejects  $\Gamma$  otherwise.

We first need

<u>Proposition 1.</u> For every integer r>0, there is a finite state acceptor  $M_r$  such that the cellular d-graph acceptor  $(\Gamma, M_r, H)$  with distinguished node D accepts all d-graphs  $\Gamma$  whose nodes are all within distance r from D in 2r+1 steps, and when it accepts, every node is in a different state.

<u>Proof</u>: Given any d-graph  $\Gamma$ , the cellular d-graph acceptor  $(\Gamma, M_{\Gamma}, H)$  operates as follows: the distinguished node D sends out a message S which propagates to the nodes at distance r from D. The paths traveled by S define a spanning tree of  $\Gamma$ , and each node is identified uniquely by marking each node's state with a sequence of arc end numbers which define the unique path from D to the node. Specifically, when a neighbor of D receives S, its state is marked with the number i if it is the ith neighbor of D. It then sends the message (S,i) to its neighbors. When an unmarked node m receives the message  $(S,i_1,\ldots,i_k)$ ,  $k\geq 1$ , from node n, and m is the jth neighbor of n, then m marks its state with  $(i_1,\ldots,i_k,j)$  and sends  $(S,i_1,\ldots,i_k,j)$  to its neighbors. If a node receives a message from more than one neighbor

simultaneously, it can choose to accept one of them, say the one sent by the lowest-numbered neighbor. Since the paths from D to each node are all different, the sequences of numbers in the states of the nodes are distinct.

If a node  $m_1$  is marked with  $(i_1, i_2, \ldots, i_r)$  and one of its neighbors, say  $m_2$ , is still unmarked, then  $m_2$  is at distance r+l away from D. A rejection signal is thus sent to D because the graph contains nodes more than distance r away. If no rejection signal is received after 2r+l steps,  $\Gamma$  is accepted.//

Given a node-labelled graph  $\alpha$  of degree  $\leq$  d, we can find its diameter r and construct its spanning tree  $T_{\alpha}$  using the method in Section 1.3.1 of [2]. The height of the spanning tree is  $\leq$  r and associated with each node is a level number. The level numbers of a node and its neighbors differ by at most 1; this follows from the way the tree is constructed. Now we can prove

<u>Proposition 2</u>. For any labelled graph  $\alpha$  of degree  $\leq$  d, there exists an  $M_{\alpha}$  such that the cellular d-graph acceptor  $(\Gamma, M, H)$  with distinguished node D accepts  $\Gamma$  if  $\alpha \simeq \Gamma$  and rejects  $\Gamma$  otherwise.

<u>Proof:</u>  $M_{\alpha}$  first simulates the action of  $M_{r}$ , where r is the diameter of  $\alpha$ , in the first r steps. It sends a rejection signal to the distinguished node D if it finds a node at distance more than r from D, since in this case  $\Gamma$  cannot be isomorphic to  $\alpha$ . At the end of step r, every node of  $\Gamma$  has

a unique identity represented in its state.

Step r+l+i (0sish) identifies the nodes corresponding to level k-i nodes of  $T_\alpha$  based only on the knowledge that the node has the right neighbors to serve as its sons in  $T_\alpha$ . In the states of these nodes, the numbers of the nodes that are to be its descendents are recorded. Thus at the end of step r+l+h, the nodes indicating that they can be the root of  $T_\alpha$  are saying that the assignments in their states are sure that all the arcs of  $T_\alpha$  exist, but the arcs in  $\alpha$  and not in  $T_\alpha$  will have to be checked. More specifically,

Step r+1: Each node decides if it can be a level h node of  $T_{\alpha}$  by looking at its label. A node that is qualified indicates this fact by recording in its state  $([n_1, \mathrm{id})], \ldots, [(n_k, \mathrm{id})])$ ,  $n_i \neq n_j$  if  $i \neq j$ , where id is the unique identity of the node obtained in previous steps, and the  $n_i$ 's are the possible level h nodes it can be. The number k is at most the number of level h nodes of  $T_{\alpha}$ , so the length of the state is bounded. All other nodes are in some neutral state.

Step r+2: Each node looks at its neighbors. If it has the right level h neighbors to serve as its sons in  $T_{\alpha}$  and qualify it to be a level h-l node, then it changes its state to  $([(\dot{n}_{1},id),\ (\dot{n}_{11},\ \dot{id}_{11}),\ldots,(\dot{n}_{1i_{1}},\ \dot{id}_{1i_{1}})],$ 

$$\begin{split} & [\,(n_2, \, \mathrm{id})\,, \,\, (n_{21}, \, \mathrm{id}_{21})\,, \ldots, (n_{2i_2}, \, \mathrm{id}_{2i_2})\,]\,, \\ & \vdots \\ & [\,(n_k, \, \mathrm{id})\,, \,\, (n_{k1}, \, \mathrm{id}_{k1})\,, \ldots, (n_{ki_k}, \, \mathrm{id}_{ki_k})\,]\,) \end{split}$$

Here  $id \neq id_{ij} \forall_{i,j}$  and  $id_{ij} \neq id_{ij}$ , if  $j \neq j'$ ; id is the unique identity of the node. Each  $[(n_j, id), \ldots, (n_{ji_j}, id_{ji_j})]$  indicates that this node can be node  $n_j$  of  $T_{\alpha}$ , the subtree of  $T_{\alpha}$  at  $n_j$  consists of nodes  $n_{j1}, \ldots, n_{ji_j}$ , and they correspond to nodes with identities  $id_{j1}, \ldots, id_{ji_j}$ , respectively. For each node, the numbers of level h-1 nodes it can be is bounded and the possible assignment from its neighbors to the subtree of  $T_{\alpha}$  is also bounded. Therefore the length of the states is bounded. All other nodes are in some neutral state regardless of their previous states.

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Step r+l+i: The qualified level h-i nodes record in their states the assignment of nodes to serve as the subtree of  $T_{\alpha}$  at the level h-i node it qualifies to be. Of course an assignment is made only if no node in the assignment corresponds to two different nodes of  $\alpha$ .

:

Step r+l+h: The possible root nodes with the assignments of all the nodes of  $\mathbf{T}_\alpha$  are known. These assignments are based on the knowledge that the arcs in  $\mathbf{T}_\alpha$  exist. Therefore each of these assignments gives a subgraph of  $\Gamma$  isomorphic to  $\mathbf{T}_\alpha$ .

Starting at Step r+h+2, each qualified root node initiates signals to check each assignment recorded in its state to make sure that all the arcs in  $\alpha$  exist and no other arcs are present. This is done by transmitting the assignment to each node. If a node not in the assignment receives the assignment signal, this means that  $\Gamma$  has more nodes than  $\alpha$  and it cannot be isomorphic to  $\alpha$ ; thus a rejection signal is sent to the distinguished node to reject  $\Gamma$ . If a node is in the assignment, when it receives the signal, it makes sure that all the arcs incident upon it are connected to the nodes with the correct identities as in  $\alpha$ . Any time a node finds an arc out of order, it sends a cancellation signal to report to the root node of this assignment to delete the assignment. If after 2h steps, the qualified root node finds that it still has uncancelled assignments, then it can send a success signal to the distinguished node D. When D gets a success signal, it accepts. However, if at the end of step r+1+h+2h+r no success signal is received by D, this means there is no successful assignment. This is because either the assignments made at step r+1+h are all cancelled or there is no assignment at all at step r+1+h, since there may be too few nodes in  $\Gamma$ , or it is not possible even to find a subgraph of  $\Gamma$  isomorphic to  $\mathbf{T}_{\alpha}$ . In any case,  $\Gamma$  is not isomorphic to  $\alpha$  and  $\Gamma$  is rejected.

An alternate method to test for isomorphism is: first give each node a unique identity as above, but starting at step r+1, the nodes' identities are transmitted to and collected by the distinguished node D. At the same time, the number of non-# nodes in  $\Gamma$  is counted and compared with k, the number of nodes in  $\alpha$ . If these two numbers are not equal, D rejects [; otherwise D sends out k! signals, each specifying an assignment of the node identities of the nodes to the nodes of  $\alpha$ . These signals propagate from neighbor to neighbor. As in the other method, each node checks its arcs as the signals are received and sends cancellation messages if any incident arc is not as in  $\alpha$ . 2r steps after D sent out the k! messages, if there is still an assignment at D, then it accepts \( \Gamma\), otherwise it rejects. Since counting and propagation of the messages all take order diameter time, this method also detects graph isomorphism in diameter time.//

#### 3. Subgraph isomorphism

#### 3.1 k-level-colored d-graphs

In this section we consider the diameter time subgraph matching problem: Given a labelled graph  $\alpha$ , find an  $M_{\alpha}$  such that ( $\Gamma$ ,  $M_{\alpha}$ , H) accepts  $\Gamma$  iff.  $\alpha$  is isomorphic to a subgraph of  $\Gamma$  in time proportional to the diameter of  $\Gamma$ . Unlike the graph isomorphism case, the number of nodes of the d-graph  $\Gamma$  may be arbitrarily large. However, the definition of  $M_{\alpha}$  depends only on  $\alpha$  and not on  $\Gamma$ ; therefore it is not possible to give each node of  $\Gamma$  a unique identification as part of its state.

Suppose  $\alpha$  has diameter r; if a node n is part of a subgraph S isomorphic to  $\alpha$ , then all the other nodes of S must be within distance r from n. We will show that if every node of  $\Gamma$  has a different state from any node within distance r from it, then we can discover whether  $\alpha$  is isomorphic to a subraph of  $\Gamma$  in time proportional to the diameter of  $\Gamma$ .

A d-graph  $\Gamma$  will be called <u>k-level-colored</u> if the nodes of  $\Gamma$  are colored and any two nodes within distance k from each other have different colors. For any node n in a d-graph  $\Gamma$ , the number of nodes within distance k from it is at most  $c(k)=d+d(d-1)+d(d-1)^2+\cdots+d(d-1)^{k-1}$ . The number of colors needed for  $\Gamma$  to be k-level-colored is no more than l+c(k). We can assume that the color at each node is part of the label and thus becomes part of the initial state of the automaton at the node, and that the color of a node is its identity.

Proposition 3. Given a labelled graph  $\beta$  with diameter r, there is a finite state automaton  $M_{\beta}$  such that for any k-level-colored (k≥r) d-graph  $\Gamma$ , the cellular d-graph acceptor ( $\Gamma$ ,  $M_{\beta}$ , H) with a distinguished node D accepts  $\Gamma$  if  $\beta \simeq a$  subraph of  $\Gamma$ , and rejects  $\Gamma$  otherwise, in time proportional to the diameter of  $\Gamma$ .

<u>Proof</u>: Given  $\beta$  we can find its spanning tree  $T_{\beta}$ . Let h be the height of  $T_{\beta}$ .  $M_{\beta}$  works in almost the same way as  $M_{\alpha}$  in Section 1 except that the first r steps of  $M_{\alpha}$  are not necessary since each node already has an identity. In the process of identifying nodes of various levels, no assignments can be made that requires two nodes of  $\beta$  to correspond to the same identity, because nodes in the same assignment must be within distance r from each other so that only one node can have a particular identity. After h+1 steps, all the possible root nodes with the correspondence between node of  $T_{\beta}$  and the node identities are known.  $M_{\beta}$  again initiates signals to check for the existence of the arcs in  $\beta$  but not in  $\mathbf{T}_{\beta}$  as  $\mathbf{M}_{\alpha}$  did with some slight modifications. The signal corresponding to each assignment is sent and transmitted from neighbor to neighbor, but it stops propagating after r steps so that it will not reach nodes at distance more than r away. When a node in the assignment receives the signal, it makes sure that the corresponding arcs in  $\beta$  incident to it all exist and are connected to the nodes with the correct identities. It does nothing if those arcs exist, but if one of the arcs

is out of order, it sends a cancellation signal to the supposed root node to delete that assignment. Again, this signal travels only h steps. All the assignments not cancelled after 2h steps are good. Thus at that point, if a supposed root node finds that it still has an uncancelled assignment, it sends a success signal to the distinguished node D. When D get s a success signal, it accepts. However, if at the end of step  $p=h+1+2h+(radius of \Gamma centered at D)$ and no success signal is received by D, it means there is no successful assignment and no subgraph of  $\Gamma$  is isomorphic Therefore we can reject  $\Gamma$ . Note that no success signal reaches D at step p iff. no success signal reaches D after step p. Therefore at step 1, besides working in the same way as  $\mathbf{M}_{\alpha}$  ,  $\mathbf{M}_{\beta}$  also sends out a special signal F from D. A return signal R is sent back to D when F reaches the leaf nodes as in the spanning tree construction of Section 1.3.1 in [2]. It takes twice the radius of  $\Gamma$  for D to receive R from all of its neighbors. Then D waits for another 3h+1 steps; if no success signal is received, it rejects .//

It should be pointed out here that each uncancelled assignment represents a subgraph isomorphic to  $\beta$ . Moreover there are redundancies because the same subgraph may be specified by different assignments which correspond to different automorphic images of the subgraph.

#### 3.2 Trees

Proposition 4. For any k>0, there is a finite state automaton  $M_k$  such that for any d-graph  $\Gamma$  which is also a tree, in time proportional to the diameter of  $\Gamma$ , the cellular d-graph automaton ( $\Gamma$ ,  $M_k$ , H) k-level-colors the nodes of  $\Gamma$ , i.e., the states of nodes within distance k from each other are distinct.

<u>Proof</u>: Let  $\ell = \lfloor \frac{k}{2} \rfloor$  whose  $\lfloor a \rfloor =$  the largest integer sa. Each node's state will have a component of the form  $(i,i_1,...,i_l)$  where  $0 \le i \le k$ ,  $0 \le i_j \le d$   $\forall 1 \le j \le l$ , which serves as the color of the node. At the first step the distinguished node writes  $(0,0,0,\ldots,0)$  in its state and sends this message to its neighbors. When an uncolored node n receives a message  $(i,i_1,\ldots,i_0)$  from its neighbor m, node n writes in its state (i+1(modulo k+1),  $i_2, i_3, \dots, i_{\ell}, j$ ) if m is the jth neighbor of n, and n sends this message to its neighbors. If an uncolored node receives messages from more than one node simultaneously, it chooses only one of them. When a node n is colored with  $(j,j_1,\ldots,j_{\ell})$ , this says that if D is the root, n is a node at level (k+1)i+j for some  $i \ge 0$  and  $j_1, \ldots, j_\ell$  specifies a path from its ancestor at level  $(k+1)i+j-\ell$  if  $j_1 \neq 0$ . If  $j_1$  is 0, then  $j_h, j_{h+1}, \dots, j_{\ell}$  is a path from D where  $j_h$  is the first nonzero element after  $j_1$ .

Given any two nodes  $n_i$  and  $n_j$  within distance k from each other, suppose their colors are  $c(n_i) = (h_i, i_1, \dots, i_{\ell})$  and  $c(n_j) = (h_j, j_1, \dots, j_{\ell})$  respectively. If  $h_i \neq h_j$ , then  $c(n_i) \neq c(n_j)$ . If  $h_i = h_j$  then  $n_i$  and  $n_j$  are on the same

level of  $\Gamma$  with root D because the levels of  $n_i$  and  $n_j$  are  $(k+1)t_1+h_i$  and  $(k+1)t_2+h_j$  for some  $t_1,t_2$ ; their difference is  $\leq k$  (otherwise  $n_i$  is not within distance k from  $n_j$ ), and this implies  $t_1=t_2$ . Let n be the closest common ancestor of  $n_i$  and  $n_j$ , i.e., no descendent of n is an ancestor of both  $n_i$  and  $n_j$ . n is at equal distance from  $n_i$  and  $n_j$ . The distance between n and  $n_i$  or  $n_j$  is  $\leq \ell = \lfloor \frac{k}{2} \rfloor$ , since otherwise the distance between  $n_i$  and  $n_j$  is > k. Therefore  $(i_1, \ldots, i_\ell) \neq (j_1, \ldots, j_\ell)$  since one contains a path from n to  $n_i$  and the other a path from n to  $n_j$ . The time it takes for the signal from n to reach a node n equals the distance between n and n. Therefore the coloring process takes diameter  $(\Gamma)$  time.//

Combining the results of Propositions 3 and 4, we see that if a d-graph  $\Gamma$  is a tree, then we can do subgraph matching in diameter ( $\Gamma$ ) time. However, because of the structure of trees -- they have no cycles, and between any two nodes there is only one simple path we can also do subgraph matching without first coloring the nodes.

Let  $\Gamma$  be a d-graph which is also a tree. Then any labelled graph isomorphic to a subgraph of  $\Gamma$  must be a tree. Let Z be a tree with root node r and height h. In the remainder of this section, we describe a cellular d-graph automaton ( $\Gamma$ ,  $M_Z$ , H) which accepts  $\Gamma$  iff.  $Z \cong a$  subgraph of  $\Gamma$ . The action of  $M_Z$  is similar to the actions of  $M_{\alpha}$  and  $M_{\beta}$  in the previous sections. Namely, at each step i ( $1 \le i \le h + 1$ )

each node n looks at its neighbors to decide if they can correspond to the sons of a level h+1-i node m in the tree Z. If they can, n declares itself to qualify as the node m of Z. The difference is that the nodes do not have distinct identities any more; rather, we know that  $\Gamma$  is a tree. Instead of recording the assignments of nodes, each node n's state simply indicates the level h-i+1 node it can be and the arc end numbers leading to its sons. Note that a node can correspond to a few different level h-i+1 nodes with different sets of sons. In the next step, the knowledge of the sons prevents those nodes from serving both as n's father and son when n corresponds to a particular node m of Z since n's neighbor can see from n's state whether it was used as n's son to qualify n as m of Z.

<u>Claim</u>: At the end of step i (lsish+1), if a node n's state indicating that it qualifies as a level h-i+1 node m of Z and its  $i_1, i_2, \dots, i_j$ -th neighbors  $n_1, n_2, \dots, n_j$  correspond to the sons  $m_1, m_2, \dots, m_j$  of m in Z, then n is the root of a tree\* isomorphic to the subtree of Z at m.

<u>Proof:</u> When i=1, the claim is clearly true since the subtree of Z at a leaf node consists of only the node itself. If the claim is true for i-1 ( $1 \le i \le h+1$ ), then  $n_1, n_2, \ldots, n_j$  are all roots of trees isomorphic to subtrees of Z at  $m_1, \ldots, m_j$ . First note that if n qualifies as corresponding

<sup>\*</sup>This tree is an acylic subgraph of \( \Gamma \).

to m, then n qualifies to be the father of  $n_1, \ldots, n_j$ . In the trees isomorphic to subtrees of  $m_1, \ldots, m_j$ , all of the  $n_k$ 's  $(1 \le k \le j)$  do not use n as a son, since otherwise n could not be that  $n_k$ 's father. Moreover, the trees at  $n_1, n_2, \ldots, n_j$  are all disjoint since they are all subgraphs of  $\Gamma$  and  $\Gamma$  is a tree. If a node A belongs to both the trees at  $n_1$  and  $n_2$ , then there exist a sequence of nodes  $n_1, n_2, \ldots, n_k$ , A joining  $n_1$  to A, and a sequence of nodes  $n_2, n_1, \ldots, n_k$ , A joining  $n_2$  to A. This means that  $n, n_1, n_1, \ldots, n_k, n_k, \ldots, n_k, n_k$  is a cycle, a contradiction. This shows that n is the root of a tree. This tree is isomorphic to the subtree of Z at  $m_i$  this is clear from the fact that the trees at  $n_k$  are isomorphic to subtrees at  $n_k$   $(1 \le k \le j)$ .//

At the end of step h+1, all the nodes that correspond to the root node of Z are identified. These nodes just send a success message to the distinguished node D for acceptance. If after h+1+height( $\Gamma$ ) steps, no success message is received by D, it rejects. Again M<sub>Z</sub> can use the same method as M<sub>B</sub> of Proposition 3 to decide when to reject.

We have thus proved

<u>Proposition 5.</u> For any labelled tree Z of height h, there is a finite state automaton  $M_Z$  such that for any d-graph  $\Gamma$  which is a tree, the cellular d-graph automaton  $(\Gamma, M_Z, H)$  accepts  $\Gamma$  iff.  $T \simeq a$  subgraph of  $\Gamma$ , and otherwise it rejects  $\Gamma$ , in time proportional to the diameter of  $\Gamma$ .

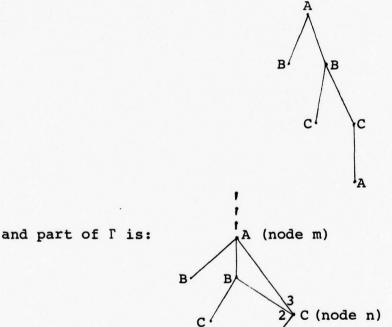
In Section 1.2.3 of [2], we showed how a cellular d-

graph automaton recognizes trees in diameter time. Combining this with the above proposition, we have the result that for any tree Z, there is a cellular d-graph automaton that recognizes all the d-graphs which are trees and have subgraphs isomorphic to Z.

#### 3.3 k-local-homogeneous d-graphs

In the last section, we exhibited a diameter time subgraph matching algorithm for trees without appealing to k-level-coloring. We have been unable to find diameter time algorithms for general d-graphs. The problems seem to be due to the existence of cycles in a general d-graph, as can be seen from the following examples:

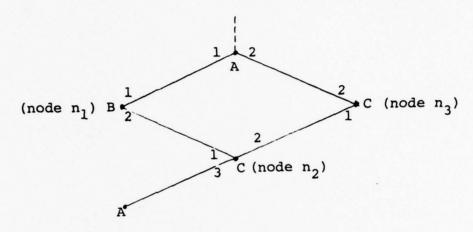
Example 1: Suppose the labelled graph  $\alpha$  is



If the states of the nodes are not different, there seems to be no way that node n can tell that its third neighbor cannot serve as its son since it is already used as its ancestor. Equivalently, there seems to be no way to prevent node m from serving as two different nodes of  $\alpha$ .

Example 2: Suppose we are looking for subgraphs to match





How can node  $n_1$  tell that node  $n_2$  is not node  $n_3$  (and thus not the desired structure) if the states of the nodes are not different? In general, how can a node distinguish the signals from different nodes? This is the same problem as in [3, 4] where the automaton cannot tell apart cycles of lengths 3 and 4.

Let us examine the special properties a tree has:

- (1) There are no nontrivial cycles in a tree. The only cycles are those consisting of a path and its exact inverse.
- (2) There is only one path between any two points.

In this section, we discuss a generalization of

these properties, and define a class of d-graphs for which diameter time subgraph matching is possible.

#### 3.3.1 Definition of k-local-homogeneity

At each node n,  $H(n) = (t_1, \dots, t_d)$  tells n that it is the  $t_i$ th neighbor of its ith neighbor. If we consider a sequence of numbers  $a_1, \dots, a_j$   $(1 \le a_i \le d \text{ for } 1 \le i \le j)$  at n as a path  $n = n_0, n_1, \dots, n_j$  such that  $n_i$  is the  $a_i$ th neighbor of  $n_{i-1}$   $(1 \le i \le j) *$ , then H(n) tells node n the inverse of any path of length 1. Define  $H^j(n) : D^j \to D^j$  such that if the image of  $(a_1, \dots, a_j)$  is  $(b_1, \dots, b_j)$  then the inverse of the path  $a_1, \dots, a_j$  is  $b_j, b_{j-1}, \dots, b_1$  and we have

$$a_1$$
  $b_1$   $a_2$   $b_2$   $a_{j-1}$   $b_{j-1}$   $a_j$   $b_j$   $a_{j-1}$   $a_{j-1}$ 

It is obvious that knowing  $H^k$  at a node n implies knowing  $H^j$  at n for any  $1 \le j \le k$ .

<u>Proposition 6</u>. For any k>0, there exists a finite state automaton  $M_k$  such that for any d-graph  $\Gamma$ , the cellular d-graph automaton ( $\Gamma$ , M, H) can find  $H^k$  and record it in its state in 2k steps.

<u>Proof:</u> Since each node knows its H-function, it can send out messages in the form of  $(i;t_i)$  to its ith neighbor  $(1 \le i \le d)$ . When a node m receives  $(i;t_i)$  from its  $t_i$ th neighbor, it

<sup>\*</sup>From now on, we will use the notations " $n=n_0,\ldots,n_j$ " and " $a_1,\ldots,a_j$  at n" interchangably.

sends out messages  $(i,j;t_i,t_i)$  to its jth neighbor  $n_i$   $(1 \le j \le d)$ where n is the  $t_{i}$  neighbor of  $n_{i}$ . In general, for  $\ell < k$ , when a node n receives  $(i_1, i_2, \dots, i_{\ell}; t_1, t_2, \dots, t_{\ell})$  from its  $t_{\ell}$ th neighbor, it augments the message and sends out  $(i_1, i_2, \dots, i_{\ell}, j; t_1, \dots, t_{\ell}, t_j)$  to its jth neighbor  $(1 \le j \le d)$ where n is the  $t_{i}^{t}$ th neighbor of its jth neighbor. If n's jth neighbor is a # node, it makes a special mark. After step k, instead of augmenting and sending out new messages, the messages backtrack, i.e., the message  $(i_1, i_2, \dots, i_k; j_1, j_2, \dots, j_k)$ travels along the path  $j_k, \dots, j_2, j_1$ . (A pointer is kept so that a message knows which  $j_{\varrho}$  to use next.) At step 2k, each node n knows from the messages it receives that the image of  $(i_1, ..., i_k)$  at n under  $H^k$  is  $(j_1, ..., j_k)$ . Hence the inverse of the path  $i_1, \dots, i_k$  from n is  $j_k, \dots, j_1$ . If a path does not reach length k because a # node is encountered, the node can tell this from the special mark at that position. Note that since each message only travels distance k away and each node has at most  $d+d(d-1)+\cdots+d(d-1)^{k-1}$  nodes within distance k from it, the number of signals at any node at each step is bounded.//

Alternate proof: At the first step, each node writes in its state the inverses of paths of length 1 from it, i.e., records its H-function. At the next step, since each node can see its neighbor's states, it can tell the inverses of paths of length 2. More specifically, suppose  $H(n) = (t_1, t_2, \ldots, t_d)$ ,  $H(m) = (s_1, s_2, \ldots, s_d)$  and m is the

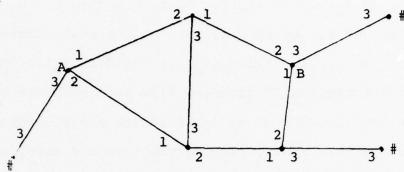
ith neighbor of n. At the end of step 1, n's state has  $((1;t_1), (2;t_2), \ldots, (d;t_d))$  and m's state has  $((1;s_1), (2;s_2), \ldots, (d;s_d))$ . If  $t_i = j$  then  $i = s_j$  by the definition of H. At step 2, part of n's state looks like this:  $(i,l;t_i,s_1)$ ,  $(i,2;t_i,s_2)$ ,..., $(i,d;t_i,s_d)$ . The same kind of information is gathered from each neighbor of n, so that the inverses of paths of length 2 are recorded in the state of n. Again, if a neighbor is a # node, a special mark is recorded in the state so that it will not attempt to extend that path further. Therefore, at each step, we can use the latest information each node acquired at the previous step to extend the paths' lengths by 1. At the end of step k, each node's state has  $H^k$  coded in it.//

In a d-graph  $\Gamma$ , a <u>node n knows all cycles of length up</u>

to k if given a sequence of arc end numbers  $a_1, \ldots, a_j$   $(1 \le a_j \le d, 1 \le j \le k)$ , hence a path of length  $j \le k$  starting from n, node n knows whether or not this path is a cycle, i.e.,  $n = n_0, n_1, \ldots, n_{j-1}, n_j = n$ , where  $n_i$  is the  $a_i$ the neighbor of  $n_{i-1}$   $(1 \le i \le j)$ . A node n knows all equivalent paths of total length up to k if given any two sequences of arc end numbers, hence two paths from n, with sum of their lengths  $\le k$ , node n knows if they lead to the same node.  $\Gamma$  is said to know all cycles of length up to k or know all equivalent paths of total length up to k iff. every node in  $\Gamma$  knows the respective information. The following example shows that knowing  $H^k$  does not imply knowing all cycles of length j nor knowing

all equivalent paths of total length j for some j>2.

#### Example:



H(n) = (2,1,3) for every node n and so  $H^k(n)$  is the same for every node. However, the cycles at node A and node B are quite different. At A, 232 gives a cycle, while at B, 232 is not a cycle. At A, 23 and 1 reach the same node, but at B, 23 and 1 do not meet.

<u>Proposition 7.</u> A node n knows all cycles of length up to k iff. n knows all equivalent paths of total length up to k.

<u>Proof</u>: For any d-graph  $\Gamma$ , there is a cellular d-graph automaton that finds  $H^k$  in constant time (depending only on k). Suppose n knows all cycles of length up to k. Given any two paths  $P_1, P_2$  whose total length  $\leq k$ , n knows  $H^k$ ; thus it can invert one of the paths, say the shorter one  $P_2$ , and append the inverse of  $P_2$  to  $P_1$ . The result is a sequence of arc end numbers of length  $\leq k$ . This sequence is a cycle iff.  $P_1$  and  $P_2$  reach the same node. But n knows all cycles of length up to k, therefore n knows all equivalent paths of total length up to k.

Conversely, suppose n knows all equivalent paths of total length up to k. Given a sequence of arc end numbers  $(a_i)$  of length  $\leq k$ , n can simply break it at some point into two, using its knowledge of  $H^k$  to find the inverse of one of them. This yields two paths at n, the sum of their lengths = length of  $(a_i) \leq k$ .  $(a_i)$  is a cycle iff. these two paths reach the same node. But n knows all the equivalent paths of total length up to k; therefore n knows all the cycles of length up to k.//

Having the equivalence given by Proposition 7, we can define a d-graph  $\Gamma$  to be <u>k-locally-homogeneous</u> if at each node n of  $\Gamma$ , H(n) and all cycles of length up to k or all equivalent paths of total length up to k are known. Clearly, a d-graph that is a tree is k-locally-homogeneous, since there are <u>no</u> cycles, and no two distinct paths can be equivalent.

#### 3.3.2 Subgraph matching problem for k-locally-homogeneous d-graphs

Let  $\omega$  be a labelled graph such that the degrees of the nodes of  $\omega$  are  $\leq$  d and the diameter of  $\omega$  is r>0\*. We will define a deterministic cellular d-graph acceptor with finite-state automaton  $M_{\omega}$  that will recognize those k-locally-homogeneous d-graphs (k>2r) having a subgraph isomorphic to  $\omega$ . For each  $\omega$ , taking any one of the nodes as the root node and using the methods described in [2], a spanning tree  $T_{\omega}$  of  $\omega$  can be constructed, where the height of  $T_{\omega}$  is h $\leq$ r.

The diameter time matching of  $\omega$  to a subgraph of any k-locally-homogeneous d-graph  $\Gamma$  will be done by first trying to identify the subgraphs of  $\Gamma$  isomorphic to  $T_{\omega}$ . These subgraphs will be stored in the state of the node corresponding to the root of  $T_{\omega}$  in the form of specifying arc end numbers leading to each node corresponding to a node of  $T_{\omega}$ . Then using homogeneity conditions, the nonspanning tree edges of  $\omega$  can be checked.

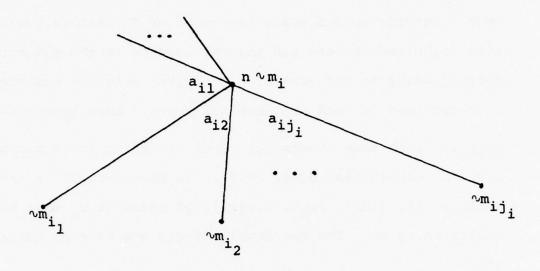
In identifying subgraphs of  $\Gamma \simeq T_{\omega}$ , the action of  $M_{\omega}$  is similar to the actions of  $M_{\alpha}$ ,  $M_{\beta}$  and  $M_{Z}$  defined earlier. Namely, at each step i (1 $\leq$ i $\leq$ h+1), each node decides if it can correspond to a node m at level h+1-i of  $T_{\omega}$  by checking if its neighbors can correspond to the sons

<sup>\*</sup>If r=0 then  $\omega$  has only one node; subgraph matching then becomes the label detecting problem of [2].

Step 2: Each node n with the proper label, and having neighbors which can correspond to the sons of a level h-l node, writes in its state:

where  $a_{it} \neq a_{it}$ , if  $t \neq t'$ . Here  $m_1, m_2, \ldots, m_s$  are the level h-1 nodes of  $T_{\omega}$  that n can correspond to, and  $m_{ij}$  are the level h nodes of  $T_{\omega}$ .  $[m_i, (a_{i1}, [m_{i1}]), \ldots, (a_{ij_i}, [m_{ij_i}])]$  means that if n corresponds to node  $m_i$ , then its  $a_{it}$ -th neighbor corresponds to node  $m_{it}$  of  $T_{\omega}$  where  $m_{it}$  (1 $\leq$ t $\leq$ j<sub>i</sub>) are the sons of  $m_i$  in  $T_{\omega}$ . This specifies a subgraph of  $\Gamma$  as shown below. Obviously, the  $m_i$ 's are not necessarily distinct. All the other non level h-1 nodes are changed to the neutral

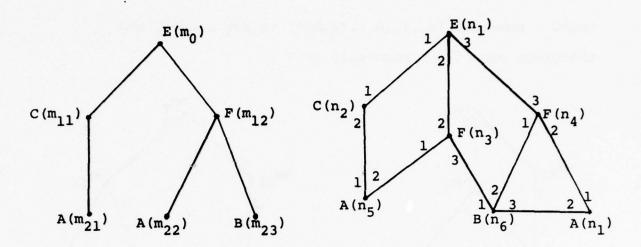
state regardless of what state they were in.



In general, at step i,  $1 \le h+1$ , a node n that qualifies to be a level h+1-i level node changes to a state of the form  $(S_1, S_2, \ldots, S_p)$  where each  $S_j$  specifies a subgraph of  $\Gamma$  at n and  $S_j = [m_j, (b_{j1}, A_{j1}), \ldots, (b_{ji_j}, A_{ji_j})]$  where  $A_j$  is  $[m_jt]$  or  $[m', (a_{p1}, A_{p1}), \ldots, (a_{p_s}, A_{p_s})]$  and  $A_j$  is in the state of the  $b_j$ -th neighbor of n.

At step h+1, since  $\mathbf{T}_{\omega}$  has only one node at level 0, all the nodes that can possibly correspond to the root of a subgraph isomorphic to  $\mathbf{T}_{\omega}$  have that subgraph recorded in their states. However, as shown by the following example, this subgraph is not necessarily a tree.

Example: Let  $T_{\omega}$  and  $\Gamma$  be the graphs shown below.

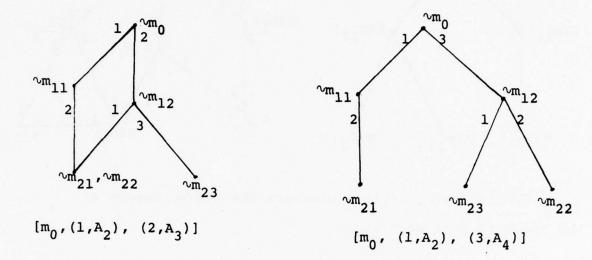


The following table summarizes the state changes of the nodes of  $\Gamma$  at each step:

Node	Step 1	Step 2	Step 3
n <sub>1</sub>	b	b	$([m_0, (1, A_2), (2, A_3)],$
n <sub>2</sub>	b	([m <sub>11</sub> ,(2,[m <sub>21</sub> ])])	$[m_0, (1, A_2), (3, A_4)])$
n <sub>3</sub>	b	$([m_{12}, (1,[m_{22}]), (3, [m_{23}])])$	b
n <sub>4</sub>	b	([m <sub>12</sub> ,(1,[m <sub>23</sub> ]),(2,[m <sub>22</sub> ])])	b
n <sub>5</sub>	$([m_{21}], [m_{22}])$	b	b
n <sub>6</sub>	([m <sub>23</sub> ])	b	b
n <sub>7</sub>	([m <sub>21</sub> ], [m <sub>23</sub> ])	b	b

where  $A_2$ ,  $A_3$ ,  $A_4$  are the states of  $n_2$ ,  $n_3$ ,  $n_4$  at step 2. The state of  $n_1$  shows that there are two subgraphs as shown

below. However,  $[m_0,(1,A_2),(2,A_3)]$  is not a tree, and therefore cannot be isomorphic to  $T_{\omega}$ .



It is easy to see (by an induction proof) that if the subgraph S obtained at a node n at step h+l is a tree, then S is isomorphic to  $\mathbf{T}_{\omega}$ . S is a tree iff. it contains no cycles. Hence, since  $\Gamma$  is k-locally-homogeneous, n can tell which subgraph in its state is not a tree, and can delete that subgraph from its state at step h+2. All the subgraphs that remain are trees isomorphic to  $\mathbf{T}_{\omega}$ . [In fact, the task of cycle checking can be done so that each node declares itself to be a level h+l-i node only if its subgraph is a tree, so that many non-tree subgraphs are deleted before step h+l.] At step h+3, node n checks to make sure the edges of  $\omega$  not in  $\mathbf{T}_{\omega}$  exist, using the knowledge it posesses about k-local homogeneity. For example, suppose  $\mathbf{m}_{\mathbf{i}}$  should

be connected to  $m_j$  in  $\omega$ , and the tree path in the subgraph from the root node n to the corresponding nodes of  $m_i$  and  $m_j$  are  $a_1, \ldots, a_{t_i}$  and  $b_1, \ldots, b_{t_j}$ ; then node n just has to make sure that there is an arc end number c such that  $a_1, \ldots, a_{t_i}$ , c and  $b_1, \ldots, b_{t_j}$  both reach  $m_j$ . If any non-tree edge does not exist, that subgraph is deleted from the state of node n.

If there is a tree left in the state of any node, then the node sends a success message to the distinguished node to signal acceptance. If the distinguished node receives no success message after step h+3+diameter( $\Gamma$ ) it rejects  $\Gamma$ . Again the distinguished node can tell that h+3+diameter( $\Gamma$ ) steps have passed by the same method used in Proposition 3.

We have thus proved

Proposition 8. For any labelled graph  $\omega$  with degree  $\leq$  d and diameter r, there exists a finite state automaton M such that for any k-locally homogeneous d-graph  $\Gamma$  (k>2r), the cellular d-graph acceptor ( $\Gamma$ , M $_{\omega}$ , H) accepts  $\Gamma$  if  $\omega \simeq$  a subgraph of  $\Gamma$ , and rejects  $\Gamma$  otherwise, in time proportional to the diameter of  $\Gamma$ .

#### 3.4 Homogeneous d-graphs

A two-dimensional array may be regarded as a 4-graph, provided we assume the boundary (#) nodes are distinct so that each # node has only one neighbor, i.e.,

node are labelled with 1(=N), 2(=W), 3(=S), 4(=E). Each node n knows the inverse of any path starting from n, since 1 and 3, 2 and 4 are always inverses of one another. Each node also knows when a path is a cycle by checking if the number of 1's = the number of 3's and the number of 2's = the number of 4's. Therefore a two-dimensional array is a k-locally-homogeneous d-graph for any  $k \ge 1$ . Moreover, all the k-local-homogeneity conditions at each node are the same in the sense that if a path from a node exists (no # node is encountered) then the same criterion determines, for all nodes, whether or not the path is a cycle.

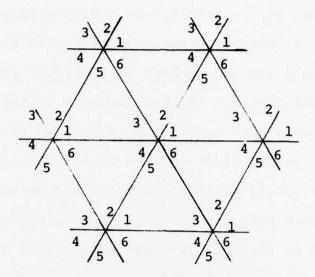
A d-graph will be called <u>k-homogeneous</u> if all the k-local-homogeneity conditions are the same for every node of Γ. If the d-graph is k-homogeneous for every k≥l, we call it simply <u>homogeneous</u>. As indicated above, the two-dimensional arrays are homogeneous 4-graphs. It is easy to see, analogously, that any n-dimensional array is a homogeneous 2n-graph.

A natural way to specify the homogeneity conditions of a d-graph is in terms of group generators and relations. We can regard the d arc end numbers at each node as the generators. A relation  $s_1s_2\cdots s_t=e$  (where e is the identity) says that at each node n, the path  $s_1s_2\cdots s_t$  is a cycle. Thus knowing the relations implies knowing all the cycles. Moreover, the cycles of length 2 at each node are the same. If  $s_1s_2=e$  then when one end of an arc is numbered with  $s_1$ , the other end of the same arc must be numbered with  $s_2$ ; thus  $s_1=s_2^{-1}$ . This shows that the d generators must form a group.

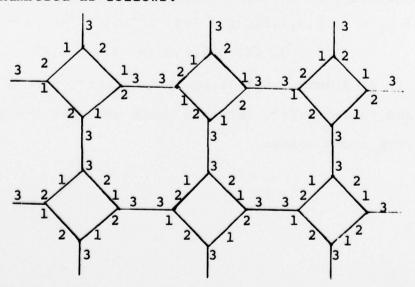
Mylopoulos and Pavlidis in [5, 6] described a number of graphs corresponding to different finitely presented abelian groups. Any finite subgraph of one of these graphs (with the appropriate # nodes added to make the degree exactly d at each non-# node) is a homogeneous d-graph. This is illustrated by the following three examples.

(1) <u>Hexagonal arrays and (semi)-regular tessellations</u>
Let  $G_1 = \{1,2,3,4,5,6\}/\{1^{-1}=4, 2^{-1}=5, 3^{-1}=6, 13=2, ij = ji \text{ for any } i,j \text{ in } 1,2,...,6\}\}$ 

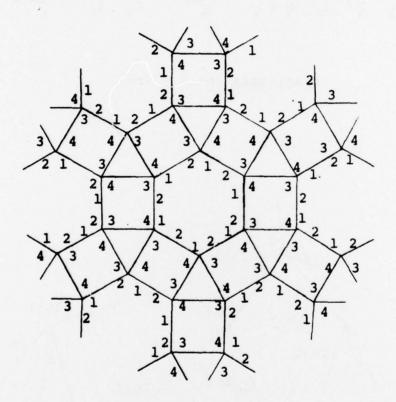
If 1, 2, 3 denote the directions "right", "above right", and "above left", then the graph of  $G_1$  is the hexagonal array shown below:



The three regular tessellations and eight semi-regular (Archimedean) tessellations in the Euclidean plane as shown on pages 24, 41 and 42 of [7] can all be regarded as homogeneous d-graphs. For example, the tessellation (4,8,8), which has three polygonal faces surrounding each vertex, where the numbers of sides of the faces are 4, 8 and 8, can be represented by the generators  $\{1,2,3\}$  and the relations  $1^{-1}=2$ ,  $3^{-1}=3$ , 1111=e, 13131313=e if the arc ends are numbered as follows:

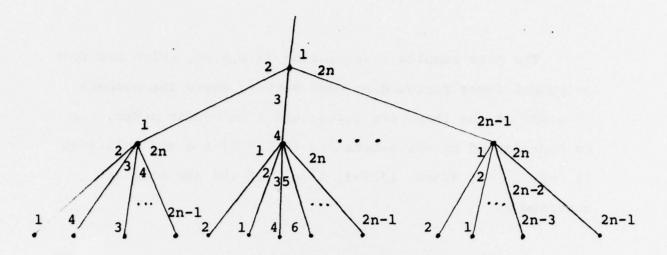


The more complex tessellation (3,4,6,4), which has four polygonal faces surrounding each vertex, where the numbers of sides of the faces are 3,4,6, and 4 in cyclic order, can be represented by the generators  $\{1,2,3,4\}$  and the relations  $\{1^{-1}=2, 3^{-1}=4, 333=e, 1313=3, 1^6=e\}$ , if the arc ends are numbered as

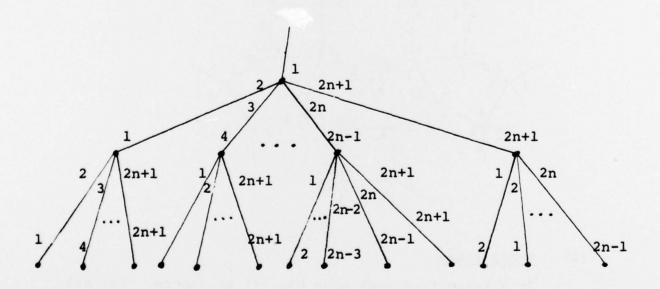


#### (2) t-ary trees

In a t-ary tree each node has t+1 neighbors. For  $n \ge 1$  let  $G_{2,n} = \{1,\ldots,2n\}/\{i^{-1}=i-1\,|\,i=2,4,\ldots,2n\}$   $G_{2,n}^{'} = \{1,\ldots,2n+1\}/(\{i^{-1}=i-1\,|\,i=2,4,\ldots,2n\}\cup\{(2n+1)^{-1}=2n+1\})\}$  Then the graph of  $G_{2,n}$  is a t-ary tree for t = 2n-1, and the graph of  $G_{2,n}^{'}$  is a t-ary tree for t = 2n.



t-ary tree for t = 2n-1



t-ary tree for t = 2n

Note that  $G_2$  and  $G_2'$  are non-abelian since 23  $\neq$  32 in both groups.

### (3) Complete graphs

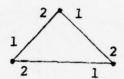
For  $n \ge 1$ , let  $D_{2n+1} = \{i^{-1} = n+i | 1 \le i \le n\} \cup \{1^{j} j = e | 2 \le j \le n\} \cup \{1^{2n+1} = e\};$ 

$$D_{2n+2} = \{i^{-1} = n+i \mid 1 \le i \le n\} \cup \{(2n+1)^{-1} = 2n+1, 1^{2n+2} = e, 1^{n+1}(2n+1) = e\}$$

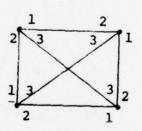
$$\cup \{1^{j} \mid 2 \le j \le n\}.$$

It is not hard to show that the graph of  $G_i = \{1, ..., i-1\}/D_i$ ,  $i \ge 3$  is  $C_i$ , the complete graph with i nodes.

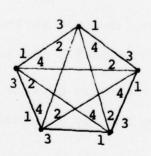
C3:



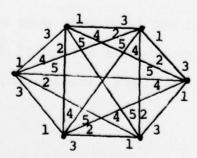
C4:



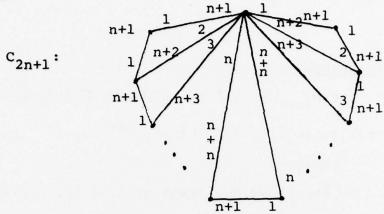
C5:

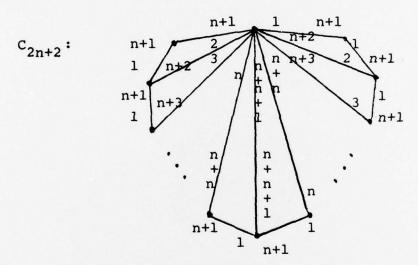


C6:



#### In general:





Since every homogeneous d-graph is k-locally-homogeneous for all k, the results of Section 2.3 imply

Proposition 9. For any homogeneous d-graph, subgraph matching can be done in diameter time by a cellular d-graph automaton.

It should be pointed out here that when we consider arrays as homogeneous d-graphs, the notion of direction in an array is not important in graph isomorphism, namely

is isomorphic to and and ... and

#### 3.5 Application: clique finding in d-graphs

In a clique all the nodes are neighbors of each other. Since every node of a d-graph has at most d non-# neighbors, the size of any clique in a d-graph is  $\leq$  d+1. This makes the clique finding problem in a d-graph much easier than that in a general graph.

Now consider the problem of finding the size of a maximal clique in a d-graph  $\Gamma$  in diameter (of  $\Gamma$ ) time. If we find the size of the largest clique at each node of  $\Gamma$ , one node at a time, and then transmit the maximum of the sizes to the distinguished node, this takes area time. But largest clique finding at many nodes simultaneously will

involve difficulties, since the signals from different nodes are not distinguishable. A better approach is to try to find subgraphs of  $\Gamma$  isomorphic to  $C_{d+1}$ , i.e., cliques of size d+1; if none exist, then we try subgraphs isomorphic to  $C_{d}$ ,  $C_{d-1}$ ,..., $C_{3}$ , $C_{2}$  in order (there are always subgraphs isomorphic to  $C_2$  if  $\Gamma$  is connected and has more than one non-# node). When for some i, a subgraph isomorphic to  $C_i$  is found, i is transmitted to the distinguished node D as the size of the maximal clique in I. When attempting to find subgraphs isomorphic to  $C_i$  (2 $\leq$ i $\leq$ d+1) in diameter time, the difficulties of subgraph matching in a general d-graph as discussed in Section 2.1 also arise. However, if  $\Gamma$  is homogeneous, or 3-locally-homogeneous or 1-level-colored, we can detect the existence or nonexistence of  $C_i$  in diameter (of Therefore the size of a maximal clique can be Γ) time. transmitted and recorded in the state of the distinguished node of  $\Gamma$  in time proportional to the diameter of  $\Gamma$ , since there are at most d C; 's to be checked.

When a subgraph isomorphic to  $C_i$  is identified, the node n of  $\Gamma$  corresponding to a special node (say A, the root of a spanning tree  $T_{C_i}$ ) of  $C_i$  can be identified and the subgraph isomorphic to  $C_i$  can be recorded in n's state. Therefore it is easy to mark the cliques of  $\Gamma$  isomorphic to  $C_i$ . Note that if the nodes of  $C_i$  have the same label, then when node n identifies itself as corresponding to node A of  $C_i$ , there are (i-1)! different correspondences of  $C_i$  to the same i-1 neighbors of n in  $\Gamma$ , since each qualifies as

any one of the i-l nodes of  $C_i$ . n can get rid of these redundant assignments by just specifying which i-l of its neighbors belong to  $C_i$ . It is also straightforward to see that each node can record in its state the size of the largest clique it belongs to and which of its neighbors form such largest cliques.

#### 4. Concluding remarks

Diameter time algorithms for the graph and subgraph matching problems are presented for trees, k-level-colored d-graphs, k-locally-homogeneous d-graphs, and homogeneous d-graphs. If fast algorithms are found to k-level-color a d-graph, then we will also have a fast algorithm for subgraph matching.

k-local-homogeneity seems somewhat artificial; however, a special case of it, namely homogeneity, holds for many important classes of d-graphs.

Homogeneous d-graphs may be considered as a natural generalization of both arrays and trees. The arc end numbering of a homogeneous d-graph is consistent. The description of the homogeneity conditions is the same at each node. In general, the description is also finite and compact (for example using group presentation), so that it can easily be stored in the finite state automaton at each node of the d-graph. It would be of interest to further study homogeneous d-graphs.

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are sets of d-graphs, i.e., labelled graphs of bounded degree whose				
arcs at each node are numbered. This report discusses acceptance tasks that depend on the concept of d-graph isomorphism in particular, the task of deciding whether a d-graph has a d-subgraph isomorphic to a				
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